Amendments to the Specification

Additions are indicated by underlining (underlining).

Please replace the paragraph on page 3 lines 16-31 with the following amended paragraph:

The present invention will now be described with reference to the following drawings, in which

- Fig. 1 shows the primary structure and disulfide bridges of the recombinant Histagged rFel d 1(2+1) fusion molecule [SEQ ID NO: 35];
- Fig. 2 shows the results of size exclusion chromatography of rFel d 1(2+1);
- Fig. 3 shows the deconvoluted electrospray mass spectrum from analysis of rFel d 1(2+1);
- Fig. 4 shows the electrospray mass spectrum from analysis of non-reduced rFel d 1(2+1) digested with trypsin;
- Fig. 5 shows the time-dependent dissociation analysis of immobilised rFel d 1(2+1) by surface plasmon resonance;
- Fig. 6 shows the result of a far-UV CD analysis of rFel d 1 and nFel d 1;
- Fig. 7 shows the IgE responses to Fel d 1 in individuals sensitised to cats in direct ELISA;
- Fig. 8 shows the dose-dependent inhibition of serum IgE from a pool of individuals sensitised to cats; and
- Fig. 9 shows the biologic activity of rFel d 1(2+1) and nFel d 1 in two patients allergic to cats.

Please replace Table 1 spanning pages 8-9 with the following amended table:

Table 1

| Primer no. | Sequence | | | |
|------------|---|--|--|--|
| 1(F) | 5'- gtacatatg g aaatctgccc ggctgttaaa cgtgacgttg acctgttcct gaccggtacc ccggacgaat acgttgaaca ggttg-3' [SEQ ID NO: 5] | | | |
| 2(R) | 5'-ggcagagett tgtactgage aacetgttea aegtattegt eegggtgage aacetgttea aegtatte-3' [SEQ ID NO: 6] | | | |
| 3(F) | 5'-tgctcagtac aaagctctgc cggttgttct ggaaaacgct cgtatcctga aaaactgcgt tgacgctaaa atgacc-3' [SEQ ID NO: 7] | | | |
| 4(R) | 5'- <u>cctctcgag</u> g cacagegggg aggtgtagat tttgtccage agggacagag egttttettt gtettetteg gteattttag egteaaege-3' [SEQ ID NO: 8] | | | |
| 5(F) | 5'- <u>qtacatatq</u> g ttaaaatggc tgaaacctgc ccgatcttct acgacgtttt cttcgctgtt gctaacggta acgaac-3' [SEQ ID NO: 9] | | | |
| 6(R) | 5'-ggtacgttcc ggttcggtag cgttaacttt ggtcagggac aggtccagca gcagttcgtt accgttagca acagc-3' [SEQ ID NO: 10] | | | |
| 7(F) | 5'-ctaccgaacc ggaacgtacc gctatgaaaa aaatccagga ctgctacgtt gaaaacggtc tgatctcccg tgttctggac-3' [SEQ ID NO: 11] | | | |
| 8(R) | 5'-gcttcaccca tgcagtcttt ggaggaggag atggtggtca taaccagacc gtccagaaca cgggagatca g-3' [SEQ ID NO: 12] | | | |
| 9(F) | 5'-caaagactgc atgggtgaag ctgttcagaa caccgttgaa gacctgaaac tgaacaccct gggtcgctcg agagg-3' [SEQ ID NO: 13] | | | |
| 10(R) | 5'- cctctcgag a cgacccaggg tg-3' [SEQ ID NO: 14] | | | |
| 11(linker) | 5'-cgtttaacag ccgggcagat ttcacgaccc agggtgttca gtttc-3' [SEQ ID NO: 15] | | | |

Please replace Table 2 spanning pages 12-13 with the following amended table:

Table 2

| Tryptic fragment | Residues | Sequence | Theoretial mass | Experimental mass |
|------------------|----------|---|-----------------|---------------------------------------|
| nagment | | | [M+H] | [M+H} |
| T5 | 44-44 | (K)K(I) [SEQ ID NO: 16] | 147.11 | not found |
| T11 | 100-100 | (K)R(D) [SEQ ID NO: 17] | 175.12 | not found |
| T14 | 132-134 | (R)ILK(N) [SEQ ID NO: 18] | 373.28 | not found |
| T1 | 0-2 | (-)MVK(M) [SEQ ID NO: 19] | 377.22 | not found |
| T4 | 40-43 | (R)TAMK(K) [SEQ ID NO: 20] | 450.24 | 450.75 |
| T9 | 87-92 | (K)LNTLGR(E) [SEQ ID NO: 21] | 673.40 | 673.40 |
| T16 | 141-146 | (K)MTEEDK(E) [SEQ ID NO: 22] | 752.31 | not found ^a |
| Т3 | 32-39 | (K)VNATEPER(T) [SEQ ID NO: 23] | 915.45 | 915.44 |
| T17 | 147-155 | (K)ENALSLLDK(I) [SEQ ID NO: 24] | 1002.55 | not found ^a |
| T13 | 122-131 | (K)ALPVVLENAR(I) [SEQ ID NO: 25] | 1081.64 | 1081/60 |
| T7 | 58-71 | (R)VLDGLVMTTISSSK(D) [SEQ ID NO: 26] | 1450.78 | 1450.72 |
| T6 | 45-57 | (K)IQD C YVENGLISR(V) [SEQ ID NO: 27] | 2156.01 | 2155.98 |
| T15 | 135-140 | (K)NCVDAK(M) [SEQ ID NO: 28] | | · · · · · · · · · · · · · · · · · · · |
| Т8 | 72-86 | (K)DCMGEAVQNTVEDLK(L) [SEQ ID NO: 29] | 2408.11 | 2408.07 |
| T10 | 93-99 | (R)EICPAVK(R) [SEQ ID NO: 30] | | |
| T12 | 101-121 | (R)DVDLFLTGTPDEYVEQVAQYK(A) [SEQ ID NO: 31] | 2430.17 | 2430.09 |
| T2 | 3-31 | (K)MAET C PIFYDVFFAVANGNELLLDLSLTK(V) [SEQ ID NO: 32] | 5092.46 | 5092.28 |
| T18 | 156-170 | (K)IYTSPLCLEHHHHHHH(-) [SEQ ID NO: 33] | | |
| T16+17 | 141-155 | (K)MTEEDKENALSLLDK(I) [SEQ ID NO: 34] | 1735.86 | 1735.80 |

^a Not found as a separate fragment. Detected as a part of the larger fragment T16 + T17 with [M+H] 1735.80, resulting from a miscleavage at Lys₁₄₆ which is surrounded by acidic residues, see Fig. 1.